

RAW SEQUENCE LISTING ERROR REPORT

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10/088,092
P41/0
1/30/2002

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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

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- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual: ePAVE)
- 2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
- 3. Hand Carry directly to:

U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202

- U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
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Revised 01/29/2002



PCT10

RAW SEQUENCE LISTING DATE: 01/30/2003 PATENT APPLICATION: US/10/088,092 TIME: 10:12:10 Input Set : A:\220738USOPCT.txt Output Set: N:\CRF4\01302003\J088092.raw 3 <110> APPLICANT: ISHIZAKI, JUN SUZUKI, NORIKO HANASAKI, KOHJI 7 <120> TITLE OF INVENTION: A GENE ENCODING NOVEL HUMAN SECRETORY TYPE PHOSPHOLIPASE A2 9 <130> FILE REFERENCE: 220738USOPCT 11 <140> CURRENT APPLICATION NUMBER: 10/088,092 C--> 12 <141> CURRENT FILING DATE: 2002-07-29 14 <150> PRIOR APPLICATION NUMBER: PCT/JP00/06344 15 <151> PRIOR FILING DATE: 2000-09-18 17 <150> PRIOR APPLICATION NUMBER: JP 11-266616 18 <151> PRIOR FILING DATE: 1999-09-21 20 <160> NUMBER OF SEQ ID NOS: 34 22 <170> SOFTWARE: PatentIn version 3.1 24 <210> SEQ ID NO: 1 **Does Not Comply** 25 <211> LENGTH: 320 Corrected Diskette Needed 26 <212> TYPE: DNA 27 <213> ORGANISM: Mus musculus 29 <220> FEATURE: "c' is at location 1 30 <221> NAME/KEY: misc_feature 31 <222> LOCATION (1)..(1) 32 <223> OTHER INFORMATION: n = a, c, g, or 35 <220> FEATURE: 36 <221> NAME/KEY: misc_feature 37 <222> LOCATION: (11)..(11) 38 <223> OTHER INFORMATION: n = a, c, g, or t 41 <220> FEATURE: 42 <221> NAME/KEY: misc_feature 43 <222> LOCATION: (29)..(29) 44 <223> OTHER INFORMATION: n = a, c, g, or t

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47 <220> FEATURE:

53 <220> FEATURE:

59 <220> FEATURE:

48 <221> NAME/KEY: misc_feature 49 <222> LOCATION: (38)..(38)

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50 < 223 > OTHER INFORMATION: n = a, c, g, or t

56 <223> OTHER INFORMATION: n = a, c, g, or t

62 <223> OTHER INFORMATION: n = a, c, g, or t



RAW SEQUENCE LISTING DATE: 01/30/2003 PATENT APPLICATION: US/10/088,092 TIME: 10:12:10

Input Set : A:\220738USOPCT.txt

W>	68 cggtggctcc cactggccag tgggacgaaa cggattggtg ttgtcatgcc catgactgct 70 gctatggccg cctggagaan ctgggctgtg accccaagct ggaaaagtac ctcttctcta 72 tcactcgaga caacatcttc tgtgctggta aaacggcttg ccagcggcat acctgcgaat	120 180 240
W>	74 gtgacaaaaa accgctctct gctttcgcca caacctgaac acttacaacc gcaantatgc	300
	76 ccactacccc aacaagctgt	320
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	85 <223> OTHER INFORMATION: Synthetic DNA	
	87 <220> FEATURE:	
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	89 <222> LOCATION: (11)(11)	
	90 <223> OTHER INFORMATION: $n = a$, c , g , or t	
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	97 <210> SEQ ID NO: 3	
	98 <211> LENGTH: 20	
	99 <212> TYPE: DNA	
	100 <213> ORGANISM: Artificial Sequence 102 <220> FEATURE:	
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	111 <212> TYPE: DNA	
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	115 <223> OTHER INFORMATION: Synthetic DNA	
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	118 <221> NAME/KEY: misc_feature	
	119 <222> LOCATION: (9)(9) 120 <223> OTHER INFORMATION: n = a, c, g, or t	
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	133 <223> OTHER INFORMATION: Synthetic DNA	
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	139 <210> SEQ ID NO: 6	
	140 <211> LENGTH: 23	
	141 <212> TYPE: DNA	
	142 <213> ORGANISM: Artificial Sequence	



RAW SEQUENCE LISTING DATE: 01/30/2003 PATENT APPLICATION: US/10/088,092 TIME: 10:12:10

Input Set : A:\220738USOPCT.txt

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		<211> LENGTH: 22	
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		<213> ORGANISM: Artificial Sequence	
		<220> FEATURE:	
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		<210> SEQ ID NO: 9	
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2	Τ.Ω	NAZON LEWING:	



RAW SEQUENCE LISTING DATE: 01/30/2003 PATENT APPLICATION: US/10/088,092 TIME: 10:12:10

Input Set: A:\220738USOPCT.txt

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     231 <400> SEQUENCE: 13
     232 agggtattga gatgccagag gc
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     242 <222> LOCATION: (166)..(591)
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     248 <222> LOCATION: (223)..()
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                                                                               120
     257 aggtggcaga gcaggetece atgeeceetg ectaeeteee eeagg atg aaa eet eee
                                                                               177
                                                            Met Lys Pro Pro
     261 att gcc ctg gct tgc ctt tgc ctc ctg gtg ccc ctg gct ggc ggg aac
                                                                               225
     262 Ile Ala Leu Ala Cys Leu Cys Leu Leu Val Pro Leu Ala Gly Gly Asn
     263 -15
                             -10
                                                  -5
                                                                  -1
     265 ctg gtc cag ttt gga gtg atg att gag aga atg acg gga aag cct gcc
                                                                               273
     266 Leu Val Gln Phe Gly Val Met Ile Glu Arg Met Thr Gly Lys Pro Ala
     267
     269 ctg cag tac aat gac tat ggc tgc tat tgc ggt gtc ggt ggc tcc cac
                                                                               321
     270 Leu Gln Tyr Asn Asp Tyr Gly Cys Tyr Cys Gly Val Gly Gly Ser His
                 20
                                     25
     273 tgg cca gtg gac gag acg gat tgg tgt tgt cat gcc cat gac tgc tgc
                                                                               369
     274 Trp Pro Val Asp Glu Thr Asp Trp Cys Cys His Ala His Asp Cys Cys
     277 tat ggc cgc ctg gag aag ctg ggc tgt gac ccc aag ctg gaa aag tac
                                                                               417
     278 Tyr Gly Arg Leu Glu Lys Leu Gly Cys Asp Pro Lys Leu Glu Lys Tyr
                             55
     281 ctc ttc tct atc act cga gac aac atc ttc tgt gct ggt aga acg gct
                                                                               465
     282 Leu Phe Ser Ile Thr Arg Asp Asn Ile Phe Cys Ala Gly Arg Thr Ala
     285 tgc cag cgg cat acc tgc gag tgt gac aag aga gct gct ctt tgc ttt
                                                                               513
     286 Cys Gln Arg His Thr Cys Glu Cys Asp Lys Arg Ala Ala Leu Cys Phe
     287
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DATE: 01/30/2003 PATENT APPLICATION: US/10/088,092 TIME: 10:12:10

Input Set : A:\220738USOPCT.txt

289 cgc cac aac ctg aac act tac aac cgc aag tat gcc cac tac ccc aac 290 Arg His Asn Leu Asn Thr Tyr Asn Arg Lys Tyr Ala His Tyr Pro Asn 291 100 105 110	561 n
293 aag ctg tgt act ggg ccc acc cca ccc tgc tgaggccctg ctcggctcca 294 Lys Leu Cys Thr Gly Pro Thr Pro Pro Cys	611
295 115 120 297 tagccacccc aggctgctgc agtctcaggc ccagagaagc tcggaaccca gattcctc	ctc 671
299 ccagcagact catecegece ecceeccaga gateatgage eetggtetet ggeeteca	agg 731
301 accacaccag atccacggga tcagctgaag aagtcacggg actcgtcagc gctcacaa	nga 791
303 tccactaagt cgcctctggc atctcaatac cctcttctga ataagagaag tctatttt	
305 cgaaaaaaaa aaaaaaaaaa aaaaaaaaaa aa	883
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316 -15 -10 -5	
319 Ala Gly Gly Asn Leu Val Gln Phe Gly Val Met Ile Glu Arg Met Thr 320 -1 1 5 10	
320 -1 1 5 10 323 Gly Lys Pro Ala Leu Gln Tyr Asn Asp Tyr Gly Cys Tyr Cys Gly Val	
324 15 20 25	e
327 Gly Gly Ser His Trp Pro Val Asp Glu Thr Asp Trp Cys Cys His Ala	L
328 30 35 40 45	
331 His Asp Cys Cys Tyr Gly Arg Leu Glu Lys Leu Gly Cys Asp Pro Lys	:
332 50 55 60	
335 Leu Glu Lys Tyr Leu Phe Ser Ile Thr Arg Asp Asn Ile Phe Cys Ala 336 70 75	
336 65 70 75 339 Gly Arg Thr Ala Cys Gln Arg His Thr Cys Glu Cys Asp Lys Arg Ala	
340 80 85 90	
343 Ala Leu Cys Phe Arg His Asn Leu Asn Thr Tyr Asn Arg Lys Tyr Ala	
344 95 100 105	
347 His Tyr Pro Asn Lys Leu Cys Thr Gly Pro Thr Pro Pro Cys	
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369 <223> OTHER INFORMATION: Synthetic DNA 371 <400> SEQUENCE: 17	
2.T /400/ 2EÃ0EMCE: I.	

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/088,092

DATE: 01/30/2003 TIME: 10:12:11

Input Set : A:\220738USOPCT.txt

Output Set: N:\CRF4\01302003\J088092.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 11,29,38,140,295 Seq#:2; N Pos. 11

Seq#:2; N Pos. 110 Seq#:4; N Pos. 29-

Seq#:18; N Pos. 197